



224378.ST25
SEQUENCE LISTING

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Deepak KUMAR
Irman AHMAD

<120> GENE SCC-112 AND DIAGNOSTIC AND THERAPEUTIC USES THEREOF

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<140> US/10/679,580
<141> 2003-10-06
<150> PCT/US02/10850
<151> 2002-04-08

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35

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 Leu Arg Ser Gln Leu Glu Leu Leu Ile Ser Pro Thr Cys Ser Cys Lys
 530 535 540
 Gln Ala Asp Ile Cys Val Arg Glu Ile Ala Arg Lys Leu Ala Asn Pro
 545 550 555 560
 Lys Gln Pro Thr Asn Pro Phe Leu Glu Met Val Lys Phe Leu Leu Glu
 565 570 575
 Arg Ile Ala Pro Val His Ile Asp Ser Glu Ala Ile Ser Ala Leu Val
 580 585 590
 Lys Leu Met Asn Lys Ser Ile Glu Gly Thr Ala Asp Asp Glu Glu Glu
 595 600 605
 Gly Val Ser Pro Asp Thr Ala Ile Arg Ser Gly Leu Glu Leu Leu Lys
 610 615 620
 Val Leu Ser Phe Thr His Pro Thr Ser Phe His Ser Ala Glu Thr Tyr
 625 630 635 640
 Glu Ser Leu Leu Gln Cys Leu Arg Met Glu Asp Asp Lys Val Ala Glu
 645 650 655
 Ala Ala Ile Gln Ile Phe Arg Asn Thr Gly His Lys Ile Glu Thr Asp
 660 665 670
 Leu Pro Gln Ile Arg Ser Thr Leu Ile Pro Ile Leu His Gln Lys Ala
 675 680 685
 Lys Arg Gly Thr Pro His Gln Ala Lys Gln Ala Val His Cys Ile His
 690 695 700
 Ala Ile Phe Thr Asn Lys Glu Val Gln Leu Ala Gln Ile Phe Glu Pro
 705 710 715 720
 Leu Ser Arg Ser Leu Asn Ala Asp Val Pro Glu Gln Leu Ile Thr Pro
 725 730 735
 Leu Val Ser Leu Gly His Ile Ser Met Leu Ala Pro Asp Gln Phe Ala
 740 745 750

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Ser Pro Met Lys Ser Val Val Ala Asn Phe Ile Val Lys Asp Leu Leu
755 760 765

Met Asn Asp Arg Ser Thr Gly Glu Lys Asn Gly Lys Leu Trp Ser Pro
770 775 780

Asp Glu Glu Val Ser Pro Glu Val Leu Ala Lys Val Gln Ala Ile Lys
785 790 795 800

Leu Leu Val Arg Trp Leu Leu Gly Met Lys Asn Asn Gln Ser Lys Ser
805 810 815

Ala Asn Ser Thr Leu Arg Leu Leu Ser Ala Met Leu Val Ser Glu Gly
820 825 830

Asp Leu Thr Glu Gln Lys Arg Ile Ser Lys Ser Asp Met Ser Arg Leu
835 840 845

Arg Leu Ala Ala Gly Ser Ala Ile Met Lys Leu Ala Gln Glu Pro Cys
850 855 860

Tyr His Glu Ile Ile Thr Pro Glu Gln Phe Gln Leu Cys Ala Leu Val
865 870 875 880

Ile Asn Asp Glu Cys Tyr Gln Val Arg Gln Ile Phe Ala Gln Lys Leu
885 890 895

His Lys Ala Leu Val Lys Leu Leu Leu Pro Leu Glu Tyr Met Ala Ile
900 905 910

Phe Ala Leu Cys Ala Lys Asp Pro Val Lys Glu Arg Arg Ala His Ala
915 920 925

Arg Gln Cys Leu Leu Lys Asn Ile Ser Ile Arg Arg Glu Tyr Ile Lys
930 935 940

Gln Asn Pro Met Ala Thr Glu Lys Leu Leu Ser Leu Leu Pro Glu Tyr
945 950 955 960

Val Val Pro Tyr Met Ile His Leu Leu Ala His Asp Pro Asp Phe Thr
965 970 975

Arg Ser Gln Asp Val Asp Gln Leu Arg Asp Ile Lys Glu Cys Leu Trp
980 985 990

Phe Met Leu Glu Val Leu Met Thr Lys Asn Glu Asn Asn Ser His Ala
995 1000 1005

Phe Met Lys Lys Met Ala Glu Asn Ile Lys Leu Thr Arg Asp Ala Gln
1010 1015 1020

Ser Pro Asp Glu Ser Lys Thr Asn Glu Lys Leu Tyr Thr Val Cys Asp
1025 1030 1035 1040

Val Ala Leu Cys Val Ile Asn Ser Lys Ser Ala Leu Cys Asn Ala Asp
1045 1050 1055

Ser Pro Lys Asp Pro Val Leu Pro Met Lys Phe Phe Thr Gln Pro Glu
1060 1065 1070

Lys Asp Phe Cys Asn Asp Lys Ser Tyr Ile Ser Glu Glu Thr Arg Val
1075 1080 1085

Leu Leu Leu Thr Gly Lys Pro Lys Pro Ala Gly Val Leu Gly Ala Val
1090 1095 1100

Asn Lys Pro Leu Ser Ala Thr Gly Arg Lys Pro Tyr Val Arg Ser Thr

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1105	1110	1115	1120
Gly Thr Glu Thr Gly Ser Asn Ile Asn Val Asn Ser Glu Leu Asn Pro			
1125	1130	1135	
Ser Thr Gly Asn Arg Ser Arg Glu Gln Ser Ser Glu Ala Ala Glu Thr			
1140	1145	1150	
Gly Val Ser Glu Asn Glu Glu Asn Pro Val Arg Ile Ile Ser Val Thr			
1155	1160	1165	
Pro Val Lys Asn Ile Asp Pro Val Lys Asn Lys Glu Ile Asn Ser Asp			
1170	1175	1180	
Gln Ala Thr Gln Gly Asn Ile Ser Ser Asp Arg Gly Lys Lys Arg Thr			
1185	1190	1195	1200
Val Thr Ala Ala Gly Ala Glu Asn Ile Gln Gln Lys Thr Asp Glu Lys			
1205	1210	1215	
Val Asp Glu Ser Gly Pro Pro Ala Pro Ser Lys Pro Arg Arg Gly Arg			
1220	1225	1230	
Arg Pro Lys Ser Glu Ser Gln Gly Asn Ala Thr Lys Asn Asp Asp Leu			
1235	1240	1245	
Asn Lys Pro Ile Asn Lys Gly Arg Lys Arg Ala Ala Val Gly Gln Glu			
1250	1255	1260	
Ser Pro Gly Gly Leu Glu Ala Gly Asn Ala Lys Ala Pro Lys Leu Gln			
1265	1270	1275	1280
Asp Leu Ala Lys Ala Ala Pro Ala Glu Arg Gln Ile Asp Leu Gln			
1285	1290	1295	

Arg

<210> 3
<211> 20
<212> PRT

<213> Artificial

<220>
<223> C-terminus of the SCC-112 Protein

<400> 3

Lys Leu Gln Asp Leu Ala Lys Lys Ala Ala Pro Ala Glu Arg Gln Ile
1 5 10 15

Asp Leu Gln Arg
20